

In the SpecificationPlease substitute the following Table 1 on page 56:

Table 1

Primers for INSP101 cloning and sequencing

Primer	Sequence (5'-3')
GCP Forward	G GGG ACA AGT TTG TAC AAA AAA GCA GGC TTC <u>GCC ACC</u> (SEQ ID NO: 11)
GCP Reverse	GGG GAC CAC TTT GTA CAA GAA AGC TGG GTT TCA ATG GTG ATG GTG ATG GTG (SEQ ID NO: 12)
INSP101-B1P-5'-F	GCA GGC TTC GCC ACC ATG GCT ACA GGC TCC CGG ACG TCC C (SEQ ID NO: 13)
INSP101-5'-R	TAG ACT CCA TTC CCC AAG AGC TTA CAA ACT CCT GGT AGG TGT CAA AGG CC (SEQ ID NO: 14)
INSP101-3'-F	AGG AGT TTG TAA GCT CTT GGG GAA TGG AGT CTA TTC CGA CAC CCT CCA ACA (SEQ ID NO: 15)
INSP101-3'-R	G TG ATG G TG ATG G TG CTA GAA GCC ACA GCT GCC CTC CAC (SEQ ID NO: 16)
INSP101-mut-F	GGG CAG CTG TGG CTT CCA CCA TCA CCA TCA CCA TTG (SEQ ID NO: 17)
INSP101-mut-R	AAT GGT GAT GGT GAT GGT GGA AGC CAC AGC TGC CCT C (SEQ ID NO: 18)
pEAK12-F	GCC AGC TTG GCA CTT GAT GT (SEQ ID NO: 19)
pEAK12-R	GAT GGA GGT GGA CGT GTC AG (SEQ ID NO: 20)
M13F	CAG GAA ACA GCT ATG ACC (SEQ ID NO: 21)
M13R	TGT AAA ACG ACG GCC AGT (SEQ ID NO: 22)
INSP101-CP1	GCT GCA ATG GCT ACA GGC TCC C (SEQ ID NO: 23)
INSP101-CP2	AGA AGC CAC AGC TGC CCT CC (SEQ ID NO: 24)

Underlined sequence = Kozak sequence**Bold** = Stop codon*Italic sequence* = His tagShaded sequence = overlap with adjacent cDNA part